

**VERSION WITH MARKINGS TO SHOW CHANGES MADE**

**In the Claims**

1. (Once Amended) An array comprising at least one pattern of probe oligonucleotide spots stably associated with the surface of a solid support, wherein each probe oligonucleotide spot of said pattern ~~corresponds to a target nucleic acid and~~ comprises an oligonucleotide probe composition made up of long oligonucleotide probes that range in length from about 50 to 120 ~~nt~~ nucleotides.

3. (Once Amended) The array according to Claim 2, wherein each probe oligonucleotide spot in said pattern ~~corresponds~~ hybridizes to a different target nucleic acid.

Cancel Claim 4.

Cancel Claim 5.

Cancel Claim 6.

10. (Once Amended) The array according to Claim 1, wherein the ~~density of spots on~~ said array ~~does~~ do not exceed a density of about 1000/cm<sup>2</sup>.

11. (Once Amended) The array according to Claim 10, wherein the ~~density of spots on~~ said array ~~does~~ do not exceed a density of about 400/cm<sup>2</sup>.

12. (Once Amended) The array according to Claim 1, wherein the ~~number of spots on~~ said array ~~ranges~~ range from about 50 to 50,000 in number.

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13. (Once Amended) The array according to Claim 1, wherein the ~~number of spots on~~ said array ~~ranges~~ range from about 50 to 10,000 in number.

14. (Once Amended) An array comprising a pattern of probe oligonucleotide spots covalently bound to the surface of a solid support, wherein each probe oligonucleotide spot ~~corresponds to a target nucleic acid and~~ comprises a long oligonucleotide probe composition made up of long oligonucleotides of from about 60 to 100 ~~nt~~ nucleotides in length, ~~wherein each of said long oligonucleotide probes exhibits substantially the same high hybridization efficiency with its respective target and low level of non-specific hybridization.~~

16. (Once Amended) The array according to Claim 15, wherein each probe oligonucleotide spot in said pattern ~~corresponds~~ hybridizes to a different target nucleic acid.

17. (Once Amended) The array according to Claim 15, wherein two or more probe oligonucleotide spots in said pattern ~~correspond~~ hybridize to the same target nucleic acid.

18. (Once Amended) The array according to Claim 14, wherein ~~the length of each of~~ said unique oligonucleotides ranges from about 65 to 90 nucleotides in length.

19. (Once Amended) The array according to Claim 14, wherein the ~~density of spots on~~ said array ~~does~~ do not exceed a density of about 1000/cm<sup>2</sup>.

20. (Once Amended) The array according to Claim 14, wherein the ~~density of spots on~~ said array ~~does~~ do not exceed a density of about 400/cm<sup>2</sup>.

21. (Once Amended) The array according to Claim 14, wherein the ~~number of spots on~~ said array ~~ranges~~ range from about 50 to 50,000 in number.

22. (Once Amended) The array according to Claim 14, wherein the ~~number of~~ spots on said array ~~ranges~~ range from about 50 to 10,000 in number.

23. (Once Amended) An array comprising a pattern of probe oligonucleotide spots of a density that does not exceed about 400 spots/cm<sup>2</sup> covalently attached to the surface of a glass support, wherein each probe oligonucleotide spot ~~corresponds to a different target nucleic acid~~ and comprises an oligonucleotide probe composition made up of long oligonucleotides of from about 65 to 90 nt nucleotides in length, ~~wherein each of said long oligonucleotides has substantially the same high hybridization efficiency for its corresponding target and the substantially the same low level of non-specific hybridization.~~

Please add the following new claims:

--36. The array according to Claim 1, wherein any variance in hybridization efficiency among any to probes of said array does not exceed about 10-fold.

37. The array according to Claim 14, wherein any variance in hybridization efficiency among any to probes of said array does not exceed about 10-fold.

38. The array according to Claim 23, wherein any variance in hybridization efficiency among any to probes of said array does not exceed about 10-fold. --

# IN THE SPECIFICATION

Please replace the paragraph beginning on page 41, line 1, with the following rewritten paragraph:

TABLE 1

Array Position	Probe Name	Probe Sequence
A1	s64_2	AC CTAGAAAGCT ATTGAGCTG GATCCGTCC TCTGATCGT AGCCCTTCT TGAAGAAATT CGGACATCTC TGCCAAAGTC TTGTGACCTG TAGCTGCCA (SEQ ID NO:3)
A2	s64_2_90	AGAAAGCTATTGAGCTGGATCCGTCCTCTGATCGTGACGCCCTTCCCTTGAAGAAATTTCCGACATCTCTGCCAAAGTCTGTGACCTGTA (SEQ ID NO:4)
A3	s64_2_80	AGCTATTTGAGCTGGATCCGTCCTCTGATCGTGACGCCCTTCCCTTGAAGAAATTTCCGACATCTCTGCCAAAGTCTTTGTGA (SEQ ID NO:5)
A4	s64_2_70	ATTGAGCTGGATCCGTCCTCTGATCGTGACGCCCTTCCCTTGAAGAAATTTCCGACATCTCTGCCAAAGTA (SEQ ID NO:6)
B1	s64_2_60	AGCTGGATCCGTCCTCTGATCGTGACGCCCTTCCCTTGAAGAAATTTCCGACATCTCTGCCA (SEQ ID NO:7)
B2	s64_2_50	AATCCGTCCTCTGATCGTGACGCCCTTCCCTTGAAGAAATTTCCGACATCTA (SEQ ID NO:8)
C1	s26_2	AAACCCAGGA AAATACCAA TCCAGATTTC TTGAAGATC TGGAAACCTTT CAGAATGACT CCTTTAGTG CTAATTGGTT GGAGCTGTGG TCCATGACCTA (SEQ ID NO:9)
C2	s26_2_90	AGGAAATACCAAATCCAGATTTC TTGAAGATCTGGAACCTTT CAGAATGACTCCTTTTAGTGCTATTGGTTGGAGCTGTGGTCCATA (SEQ ID NO:10)
C3	s26_2_80	AATACCAAATCCAGATTTC TTGAAGATCTGGAACCTTT CAGAATGACTCCTTTAGTGCTATTGGTTGGAGCTGTGGA (SEQ ID NO:11)
C4	s26_2_70	AAATCCAGATTTC TTGAAGATCTGGAACCTTT CAGAATGACTCCTTTAGTGCTATTGGTTGGAGCA (SEQ ID NO:12)
D1	s26_2_60	ACAGATTTC TTGAAGATCTGGAACCTTT CAGAATGACTCCTTTAGTGCTATTGGTTA (SEQ ID NO:13)
D2	s26_2_50	ATTCTTTGAAGATCTGGAACCTTT CAGAATGACTCCTTTAGTGCTATTA (SEQ ID NO:14)
A5 and E5	c370_2	AGGGTC AGCTGATCTA CGAGTCTGCC ATCACCTGTG AGTACCTGGA TGAAGCATAC CCAGGGAAGA AGCTGTGCC GGATGACCCC TATGAGAAAG CTTGCA (SEQ ID NO:15)
A6 and E6	c370_2_90	AAGCTGATCTACGAGTCTGCCATCACCTGTGAGTACCTGGATGAAGCATACCCAGGGAAGAAGCTGTTCCGGATGACCCCCTATGAGAAA (SEQ ID NO:16)
A7 and E7	c370_2_80	AATCTACGAGTCTGCCATCACCTGTGAGTACCTGGATGAAGCATACCCAGGGAAGAAGCTGTTCCCGATGACCCCTATA (SEQ ID NO:17)
A8 and E8	c370_2_70	ACGAGTCTGCCATCACCTGTGAGTACCTGGATGAAGCATACCCAGGGAAGAAGCTGTTCCCGATGACCA (SEQ ID NO:18)
B5 and F5	c370_2_60	ACTGCCATCACCTGTGAGTACCTGGATGAAGCATACCCAGGGAAGAAGCTGTTGCCGGAA (SEQ ID NO:19)
B6 and F6	c370_2_50	AATCACCTGTGAGTACCTGGATGAAGCATACCCAGGGAAGAAGCTGTTGA (SEQ ID NO:20)

<b>G1</b>	s91 3	AGGCCCAAT GGCTGGAAT CTCGCTATT TAGGCATTCT ACTCAGAAAA ACCTTAAAA TTCACAAATG TGTCAAGA GCCTTGATGT GGAACCGATA (SEQ ID NO:21)
<b>G2</b>	s91_3_90	ACAAATGGCTGGAATCTCGCCTATTTAGGCATTCTACTCAGAAAAACCTTAAAAATTCACAAATGTGTCAAGAGCCCTTGATGTGGAA (SEQ ID NO:22)
<b>G3</b>	s91 3 80	AGGCTGGAATCTCGCCTATTTAGGCATTCTACTCAGAAAAACCTTAAAAATTCACAAATGTGTCAAGAGCCCTTGATA (SEQ ID NO:23)
<b>G4</b>	s91 3 70	AGAAATCTCGCCTATTTAGGCATTCTACTCAGAAAAACCTTAAAAATTCACAAATGTGTCAAGAGCCCA (SEQ ID NO:24)

Please replace the paragraph beginning on page 42, line 1, with the following rewritten paragraph:

<b>H1</b>	s91 3 60	ACTCGCCTATTTAGGCATTCTACTCAGAAAAACCTTAAAAATTCACAAATGTGTCAGAAA (SEQ ID NO:25)
<b>H2</b>	s91 3 50	ACTATTTAGGCATTCTACTCAGAAAAACCTTAAAAATTCACAAATGTGTA (SEQ ID NO:26)
<b>E1</b>	s97 4	ATAGGAGGGG TGAAGCCCAAG CTGCTCATGA ACGAGTTTGA GTCAGCCCAAG GGTGACTTTG AGAAAGTGCT GGAAGTAAAC CCCAGATA AGGCTGCAAGA (SEQ ID NO:27)
<b>E2</b>	s97_4_90	AGGGTGAAGCCCAAGCTGCTCATGAACGAGTTTGAGTCAGCAAGGGTGACTTTGAGAAAGTGCTGGAAGTAAACCCCAAGATAAGGCA (SEQ ID NO:28)
<b>E3</b>	s97 4 80	AGAAAGCCCAAGCTGCTCATGAACGAGTTTGAGTCAGCAAGGGTGACTTTGAGAAAGTGCTGGAAGTAAACCCCAAGATA (SEQ ID NO:29)
<b>E4</b>	s97 4 70	ACCAGCTGCTCATGAACGAGTTTGAGTCAGCAAGGGTGACTTTGAGAAAGTGCTGGAAGTAAACCCCA (SEQ ID NO:30)
<b>F1</b>	s97 4 60	ATGCTCATGAACGAGTTTGAGTCAGCAAGGGTGACTTTGAGAAAGTGCTGGAAGTAA (SEQ ID NO:31)
<b>F2</b>	s97 4 50	AATGAACGAGTTTGAGTCAGCAAGGGTGACTTTGAGAAAGTGCTGGA (SEQ ID NO:32)
<b>C5</b>	s74 3	ATATGT AACTGAAGAA GGTGACAGTC CTTTGGGTGA CCATGTGGGT TCTCTGTCAG AGAAATTAGC AGCAGTCGTC AATAACCTAA ATACTGGCA AGTGTA (SEQ ID NO:33)
<b>C6</b>	s74_3_90	AAACTGAAGAAAGGTGACAGTCCTTTGGGTGACCATGTGGGTTCTCTGTCAAGAAATTAGCAGCAGTCGTCAATAACCTAAATACTGGGA (SEQ ID NO:34)
<b>C7</b>	s74 3 80	AAAGAAAGGTGACAGTCCTTTGGGTGACCATGTGGGTTCTCTGTCAAGAAATTAGCAGCAGTCGTCAATAACCTAAATA (SEQ ID NO:35)
<b>C8</b>	s74 3 70	AAGTGACAGTCCTTTGGGTGACCATGTGGGTTCTCTGTCAAGAAATTAGCAGCAGTCGTCAATAACCTA (SEQ ID NO:36)
<b>D5</b>	s74 3 60	ACAGTCCTTTGGGTGACCATGTGGGTTCTCTGTCAAGAAATTAGCAGCAGTCGTCAATA (SEQ ID NO:37)
<b>D6</b>	s74 3 50	ACTTTGGGTGACCATGTGGGTTCTCTGTCAAGAAATTAGCAGCAGTCGA (SEQ ID NO:38)

TABLE 1 (CONT)